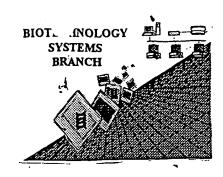
## RAW SEQUENCE LISTING ERROR REPORT



0590

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/993,756						
Source:	OIPE						
Date Processed by STIC:	12/5/01						

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

## OIPE

```
RAW SEQUENCE LISTING
                                                                DATE: 12/05/2001
                      PATENT APPLICATION: US/09/993,756
                                                                TIME: 10:42:44
                      Input Set : A:\PF0111-3CONSEQLST.txt
                      Output Set: N:\CRF3\11212001\I993756.raw
                                                                     Does Not Comply
                                                                     Corrected Diskette Needed
      1 <110> APPLICANT: Bailleul, Bernard
                                                                      Errors on P. 2
              Strosberg, A. Donny
              Akerblom, Ingrid E.
      5 <120> TITLE OF INVENTION: NOVEL HUMAN LEPTIN RECEPTOR GENE-RELATED PROTEIN
      7 <130> FILE REFERENCE: PF-0111-3 CON
     9 <140> CURRENT APPLICATION NUMBER: US/09/993,756
C--> 10 <141> CURRENT FILING DATE: 2001-11-05
     12 <160> NUMBER OF SEQ ID NOS: 4
     13 <170> SOFTWARE: PERL Program
     15 <210> SEQ ID NO: 1
     16 <211> LENGTH: 131
     17 <212> TYPE: PRT
     18 <213> ORGANISM: Homo sapiens
     20 <220> FEATURE:
     21 <221> NAME/KEY: misc_feature
     22 <223> OTHER INFORMATION: Incyte ID No: 492703
     24 <400> SEQUENCE: 1
     25 Met Ala Gly Val Lys Ala Leu Val Ala Leu Ser Phe Ser Gly Ala
                           5
     27 Ile Gly Leu Thr Phe Leu Met Leu Gly Cys Ala Leu Glu Asp Tyr
                          20
     29 Gly Val Tyr Trp Pro Leu Phe Val Leu Ile Phe His Gly Ile Ser
                                                                    45
                          35
                                               40
     31 Pro Ile Pro His Phe Ile Ala Lys Arg Val Thr Tyr Asp Ser Asp
     32
                          50
                                               55
                                                                    60
     33 Ala Thr Ser Ser Ala Cys Arg Glu Leu Ala Tyr Phe Phe Thr Thr
                          65
     35 Gly Ile Val Val Ser Ala Phe Gly Phe Pro Val Ile Leu Ala Arg
                          80
                                               85
     37 Val Ala Val Ile Lys Trp Gly Ala Cys Gly Leu Val Leu Ala Gly
                          95
                                              100
     39 Asn Ala Val Ile Phe Leu Thr Ile Gln Gly Phe Phe Leu Ile Phe
                         110
                                              115
                                                                   120
     41 Gly Arg Gly Asp Asp Phe Ser Trp Glu Gln Trp
                         125
     44 <210> SEO ID NO: 2
     45 <211> LENGTH: 874
     46 <212> TYPE: DNA
     47 <213> ORGANISM: Homo sapiens
     49 <220> FEATURE:
     50 <221> NAME/KEY: misc_feature
     51 <223> OTHER INFORMATION: Incyte ID No: 492703
     53 <220> FEATURE:
                                                                This is normally used to
Captain location of no and what
residue no represents
     54 <221> NAME/KEY: unsure
     55 <222> LOCATION: 40, 852, 862
     56 <223> OTHER INFORMATION: (a, t, c, g, or other
     58 <400> SEQUENCE: 2
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RAW SEQUENCE LISTING
                                                              DATE: 12/05/2001
                     PATENT APPLICATION: US/09/993,756
                                                              TIME: 10:42:44
                     Input Set : A:\PF0111-3CONSEQLST.txt
                     Output Set: N:\CRF3\11212001\1993756.raw
     59 gtctggcttg ggcaggctgc ccgggccgtg gcaggaagqs ggaagcagcc gcggccccag 60
     60 ttcgggagac atggcgggcg ttaaagctct cgtggcatta tccttcagtg gggctattgg 120
     61 actgactttt cttatgctgg gatgtgcctt agaggattat ggcgtttact ggcccttatt 180
     62 cgtcctgatt ttccacggca tctcccccat cccccatttc attgccaaaa gagtcaccta 240
     63 tgactcagat gcaaccagta gtgcctgtcg ggaactggca tatttcttca ctactggaat 300
     64 tgttgtttct gcctttggat ttcctgttat tcttgctcgt gtggctgtga tcaaatgggg 360
     65 agectgegge ettgtgttgg eaggeaatge agteatttte ettaeaatte aagggttttt 420
     66 ccttatattt ggaagaggag atgattttag ctgggagcag tggtagcact ttattctgat 480
     67 tacagtgcat tgaatttett agaacteata etatetgtat acatgtgcae atgeggeatt 540
     68 ttactatgaa atttaatatg ctgggttttt taataccttt atatatcatg ttcactttaa 600
     69 gaaagacttc ataagtagga gatgagtttt attctcagca aatagacctg tcaaatttag 660
     70 attatgttac tcaaattatg ttacttgttt ggctgttcat gtagtcacgg tgctctcaga 720
     71 aaatatatta acgcagtett gtaggeaget geeacettat geagtgeate gaaacetttt 780
     72 gcttqqqqat gtgcttqqaq aqqcaqataa cqctqaaqca gqcctctcat gacccaggaa 840
     73 ggccggggtg g@tccctctt t(k)ttttgtag tcca
                                                                            874
     75 <210> SEQ ID NO: 3
                                  - w can only represent a crt/u
                                                                            - k can only represent
gurtly
     76 <211> LENGTH: 145
     77 <212> TYPE: PRT
     78 <213> ORGANISM: Homo sapiens
     80 <300> PUBLICATION INFORMATION:
     81 <308> DATABASE ACCESSION NO: Genbank ID No: g733888
                                                -> 2400 > Should be used
W--> 83 <300> PUBLICATION INFORMATION:) 3
     84 Met Cys Cys His Ile His Ile Gln Cys Phe Asp Cys Cys Ser Met
                                              10
                                                                  15
     86 Lys Asn Thr Ile Leu Ala Val Ala Ala Leu Ala Phe Ala Gly Val
                         20
                                              25
                                                                  30
     88 Val Gly Leu Thr Phe Leu Val Leu Gly Cys Ala Leu Pro Arg Tyr
                         35
                                                                   45
     90 Gly Thr Trp Thr Pro Met Phe Val Ile Thr Phe Tyr Val Leu Ser
                         50
     92 Pro Val Pro Leu Leu Ile Ala Arg Arg Phe Gln Glu Asp Met Thr
                                                                  75
                                              70
                         65
     94 Gly Thr Asn Ala Cys Ile Glu Leu Ala Leu Phe Ile Thr Thr Gly
                                                                  90
                         80
                                              85
     96 Ile Val Ile Ser Ala Phe Ala Leu Pro Ile Val Leu Ala His Ala
                         95
                                             100
     98 Gly Thr Ile Ala Met Ser Ala Cys Phe Leu Ile Phe Ile Ala Asn
     99
                        110
                                             115
     100 Ser Ile Asn Phe Ser Val Ile Ile Phe Tyr Phe Arg Ile Phe Asn
                         125
                                              130
                                                                  135
     102 Gly Glu Asp Met Asn Gly Met Ser Leu Trp
     103
                         140
                                              145
     105 <210> SEQ ID NO: 4
     106 <211> LENGTH: 140
     107 <212> TYPE: PRT
     108 <213> ORGANISM: Homo sapiens
     110 <300> PUBLICATION INFORMATION:
     111 <308> DATABASE ACCESSION NO: Genbank ID No: g1197072
W--> 113 ($300) PUBLICATION INFORMATION: 4
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⇒<400>

RAW SEQUENCE LISTING DATE: 12/05/2001 PATENT APPLICATION: US/09/993,756 TIME: 10:42:44

Input Set : A:\PF0111-3CONSEQLST.txt
Output Set: N:\CRF3\11212001\1993756.raw

114 115	Met 1	Met	Glu	Phe	Lys 5	Val	Ser	Pro	Leu	Thr 10	Lys	Ile	Ile	Ser	Leu 15
	_	Gly	Phe	Leu		Leu	Gly	Phe	Leu		Val	Ile	Leu	Ser	
	Ala	Leu	Phe	His		Tyr	Tyr	Pro	Leu		Asp	Ile	Leu	Ile	
	Leu	Leu	Ala	Pro		Pro	Asn	Thr	Ile		Asn	Ala	Gly	Asn	
	Tyr	His	Thr	Ser		Phe	Met	Ser	Asp	_	Ser	Asn	Thr	Gly	
	Asp	Leu	Ala	His		Leu	Thr	Gly	Met		Val	Thr	Ser	Gly	
	Ala	Leu	Pro	Val		Phe	Tyr	His	Cys		Leu	Ile	Gly	His	
	Ser	Суѕ	Ile	Met		Met	Ile	Gly	Gly		Ile	Ile	Tyr	Ser	
	Ile	Val	Ile	Phe		Trp	Phe	Phe	Lys		Asp	Phe	Asn	Glu	
	Asp	Ser	Leu	Phe						130					133

VERIFICATION SUMMARYDATE: 12/05/2001PATENT APPLICATION: US/09/993,756TIME: 10:42:45

Input Set : A:\PF0111-3CONSEQLST.txt
Output Set: N:\CRF3\11212001\1993756.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:83 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:3 L:113 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:4